



RAW SEQUENCE LISTING  
ERROR REPORT

HG

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/613,591A  
Source: 1647  
Date Processed by STIC: 3-9-01

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

**Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is 32K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be downloaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

RECEIVED  
MAY 22 2001  
U.S. PATENT AND TRADEMARK OFFICE  
U.S. GOVERNMENT PRINTING OFFICE: 1999 50-133-100  
2001 50-133-100

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/613,591A

DATE: 03/09/2001

TIME: 11:17:20

Input Set : A:\A-378CIP5 (US).txt  
Output Set: N:\CRF3\03092001\I613591A.raw*Does Not Comply  
Corrected Diskette Needed*

3 <110> APPLICANT: BOYLE, WILLIAM  
4 LACEY, DAVID  
5 CALZONE, FRANK  
6 CHANG, MING-SHI  
7 SENALDI, GIORGIO  
9 <120> TITLE OF INVENTION: COMBINATION THERAPY FOR CONDITIONS LEADING TO BONE LOSS  
11 <130> FILE REFERENCE: A-378CIP5  
13 <140> CURRENT APPLICATION NUMBER: US 09/613,591A  
14 <141> CURRENT FILING DATE: 2000-07-10  
16 <150> PRIOR APPLICATION NUMBER: US 09/457,647  
17 <151> PRIOR FILING DATE: 1999-12-09  
19 <150> PRIOR APPLICATION NUMBER: US 09/350,670  
20 <151> PRIOR FILING DATE: 1999-07-09  
22 <150> PRIOR APPLICATION NUMBER: US 08/706,945  
23 <151> PRIOR FILING DATE: 1996-09-03  
25 <150> PRIOR APPLICATION NUMBER: US 08/577,788  
26 <151> PRIOR FILING DATE: 1995-12-22  
28 <160> NUMBER OF SEQ ID NOS: 168  
30 <170> SOFTWARE: PatentIn version 3.0  
32 <210> SEQ ID NO: 1  
33 <211> LENGTH: 36  
34 <212> TYPE: DNA  
35 <213> ORGANISM: Artificial Sequence  
37 <220> FEATURE:  
38 <221> NAME/KEY: misc\_feature  
39 <222> LOCATION: (16)..(23)  
40 <223> OTHER INFORMATION: NotI restriction site of the artificial sequence.  
43 <220> FEATURE:  
44 <221> NAME/KEY: misc\_feature  
45 <222> LOCATION: (28)..(35)  
46 <223> OTHER INFORMATION: N = random nucleic acid  
49 <400> SEQUENCE: 1  
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53 <210> SEQ ID NO: 2  
54 <211> LENGTH: 16  
55 <212> TYPE: DNA  
56 <213> ORGANISM: Artificial Sequence  
58 <220> FEATURE:  
59 <221> NAME/KEY: misc\_feature  
60 <223> OTHER INFORMATION: ds oligonucleotide adapter 16  
63 <400> SEQUENCE: 2  
64 tcgacccacg cgtccg  
67 <210> SEQ ID NO: 3  
68 <211> LENGTH: 12  
69 <212> TYPE: DNA  
70 <213> ORGANISM: Artificial Sequence  
72 <220> FEATURE:

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73 <221> NAME/KEY: misc\_feature  
74 <223> OTHER INFORMATION: ds oligonucleotide adapter  
77 <400> SEQUENCE: 3  
78 gggtgcgca 12  
81 <210> SEQ ID NO: 4  
82 <211> LENGTH: 18  
83 <212> TYPE: DNA  
84 <213> ORGANISM: Artificial Sequence  
86 <220> FEATURE:  
87 <221> NAME/KEY: misc\_feature  
88 <223> OTHER INFORMATION: PCR primer  
91 <400> SEQUENCE: 4  
92 tgtaaaacga cggccagt 18  
95 <210> SEQ ID NO: 5  
96 <211> LENGTH: 18  
97 <212> TYPE: DNA  
98 <213> ORGANISM: Artificial Sequence  
100 <220> FEATURE:  
101 <221> NAME/KEY: misc\_feature  
102 <223> OTHER INFORMATION: PCR primer  
105 <400> SEQUENCE: 5  
106 cagggaaacag ctatgacc 18  
109 <210> SEQ ID NO: 6  
110 <211> LENGTH: 20  
111 <212> TYPE: DNA  
112 <213> ORGANISM: Artificial Sequence  
114 <220> FEATURE:  
115 <221> NAME/KEY: misc\_feature  
116 <223> OTHER INFORMATION: T3 primer  
119 <400> SEQUENCE: 6  
120 caatttaaccc tcactaaagg 20  
123 <210> SEQ ID NO: 7  
124 <211> LENGTH: 23  
125 <212> TYPE: DNA  
126 <213> ORGANISM: Rattus rattus  
128 <400> SEQUENCE: 7  
129 gcatttatgac ccagaaaaccg gac 23  
132 <210> SEQ ID NO: 8  
133 <211> LENGTH: 23  
134 <212> TYPE: DNA  
135 <213> ORGANISM: Rattus rattus  
137 <400> SEQUENCE: 8  
138 aggttagcgcc cttcctcaca ttc 23  
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142 <211> LENGTH: 30  
143 <212> TYPE: DNA  
144 <213> ORGANISM: Artificial Sequence  
146 <220> FEATURE:  
147 <221> NAME/KEY: misc\_feature

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148 <223> OTHER INFORMATION: PCR primer
151 <400> SEQUENCE: 9
152 gactagtcggccaaatgaaca agtggctgtg 30
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156 <211> LENGTH: 45
157 <212> TYPE: DNA
158 <213> ORGANISM: Artificial Sequence
160 <220> FEATURE:
161 <221> NAME/KEY: misc_feature
162 <223> OTHER INFORMATION: PCR primer
165 <400> SEQUENCE: 10
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171 <212> TYPE: DNA
172 <213> ORGANISM: Artificial Sequence
174 <220> FEATURE:
175 <221> NAME/KEY: misc_feature
176 <223> OTHER INFORMATION: PCR primer
179 <400> SEQUENCE: 11
180 gcctcttagaa agagctggga c 21
183 <210> SEQ ID NO: 12
184 <211> LENGTH: 21
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:
189 <221> NAME/KEY: misc_feature
190 <223> OTHER INFORMATION: PCR primer
193 <400> SEQUENCE: 12
194 cgccgtgttc catttatgag c 21
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198 <211> LENGTH: 24
199 <212> TYPE: DNA
200 <213> ORGANISM: Rattus rattus
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203 atcaaaggca gggcatactt cctg 24
206 <210> SEQ ID NO: 14
207 <211> LENGTH: 24
208 <212> TYPE: DNA
209 <213> ORGANISM: Rattus rattus
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212 gttgcactcc tgtttcacgg tctg 24
215 <210> SEQ ID NO: 15
216 <211> LENGTH: 24
217 <212> TYPE: DNA
218 <213> ORGANISM: Rattus rattus
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221 caagacaccc tgaaggcgt gatg 24
224 <210> SEQ ID NO: 16

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225 <211> LENGTH: 24	
226 <212> TYPE: DNA	
227 <213> ORGANISM: Rattus rattus	
229 <400> SEQUENCE: 16	
230 taactttac agaaga <del>g</del> cat cagc	24
233 <210> SEQ ID NO: 17	
234 <211> LENGTH: 33	
235 <212> TYPE: DNA	
236 <213> ORGANISM: Rattus rattus	
238 <400> SEQUENCE: 17	
239 agcgccggccg catgaacaag tggctgtgct gcg	33
242 <210> SEQ ID NO: 18	
243 <211> LENGTH: 31	
244 <212> TYPE: DNA	
245 <213> ORGANISM: Rattus rattus	
247 <400> SEQUENCE: 18	
248 agtcttagag aaacagccca gtgaccatTC c	31
251 <210> SEQ ID NO: 19	
252 <211> LENGTH: 24	
253 <212> TYPE: DNA	
254 <213> ORGANISM: Rattus rattus	
256 <400> SEQUENCE: 19	
257 gtgaagctgt gcaagaacct gatg	24
260 <210> SEQ ID NO: 20	
261 <211> LENGTH: 24	
262 <212> TYPE: DNA	
263 <213> ORGANISM: Rattus rattus	
265 <400> SEQUENCE: 20	
266 atcaaaggca gggcataCTT cctg	24
269 <210> SEQ ID NO: 21	
270 <211> LENGTH: 24	
271 <212> TYPE: DNA	
272 <213> ORGANISM: Homo sapiens	
274 <400> SEQUENCE: 21	
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278 <210> SEQ ID NO: 22	
279 <211> LENGTH: 33	
280 <212> TYPE: DNA	
281 <213> ORGANISM: Homo sapiens	
283 <400> SEQUENCE: 22	
284 agcgccggccg cggggaccac aatgaacaag ttG	33
287 <210> SEQ ID NO: 23	
288 <211> LENGTH: 33	
289 <212> TYPE: DNA	
290 <213> ORGANISM: Homo sapiens	
292 <400> SEQUENCE: 23	
293 agtcttagaa ttgtgaggaa acagctcaat ggc	33
296 <210> SEQ ID NO: 24	
297 <211> LENGTH: 39	

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Input Set : A:\A-378CIP5 (US).txt  
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298 <212> TYPE: DNA
299 <213> ORGANISM: Artificial Sequence
301 <220> FEATURE:
302 <221> NAME/KEY: misc_feature
303 <223> OTHER INFORMATION: PCR primer
306 <400> SEQUENCE: 24
307 atagcgcccg ctgagccaa atcttgtac aaaactcac 39
310 <210> SEQ ID NO: 25
311 <211> LENGTH: 45
312 <212> TYPE: DNA
313 <213> ORGANISM: Artificial Sequence
315 <220> FEATURE:
316 <221> NAME/KEY: misc_feature
317 <223> OTHER INFORMATION: PCR primer
320 <400> SEQUENCE: 25
321 tctagatcg acttattcatt taccggaga cagggagagg ctctt 45
324 <210> SEQ ID NO: 26
325 <211> LENGTH: 38
326 <212> TYPE: DNA
327 <213> ORGANISM: Mus musculus
329 <400> SEQUENCE: 26
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333 <210> SEQ ID NO: 27
334 <211> LENGTH: 43
335 <212> TYPE: DNA
336 <213> ORGANISM: Mus musculus
338 <400> SEQUENCE: 27
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342 <210> SEQ ID NO: 28
343 <211> LENGTH: 38
344 <212> TYPE: DNA
345 <213> ORGANISM: Mus musculus
347 <400> SEQUENCE: 28
348 cctctgagct caagcttccg aggaccacaa tgaacaag 38
351 <210> SEQ ID NO: 29
352 <211> LENGTH: 24
353 <212> TYPE: DNA
354 <213> ORGANISM: Homo sapiens
356 <400> SEQUENCE: 29
357 tccgttaagaa acagccccgt gacc 24
360 <210> SEQ ID NO: 30
361 <211> LENGTH: 31
362 <212> TYPE: DNA
363 <213> ORGANISM: Mus musculus
365 <400> SEQUENCE: 30
366 cctctgcggc cgctgttgc tttccttct 31
369 <210> SEQ ID NO: 31
370 <211> LENGTH: 19
371 <212> TYPE: PRT

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/613,591A

DATE: 03/09/2001

TIME: 11:17:22

Input Set : A:\A-378CIP5 (US).txt

Output Set: N:\CRF3\03092001\I613591A.raw

L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1